****
Reviewer Comments:
1.
SEQUENCE LISTING
<110 >Tonen Corporation
<120 >Method for Detection or Measurement of Hepatitis C V irus
<160 >8
<210 >1
<211 >177
<212 >PRT
<213 >Hepatitis C virus
<400 >1
Met Lys Ala Ile Phe Val Leu Lys Gly Ser Leu Asp Arg Asp Pro Glu
5 10 15
Phe Met Gly Thr Asn Pro Lys Pro Gin Arg Lys Thr Lys Arg Asn Thr
 20 25 30
Asn Arg Arg Pro Gin Asp Val Lys Phe Pro Gly Gly Gly Gin Ile Val
 35 40 45
Gly Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg
 50 55 60
Ala Thr Arg Lys Thr Ser Lys Arg Ser Gin Pro Arg Gly Gly Arg Arg
 65 70 75 80
Pro Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro
85 90 95
Gly Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly
 100 105 110
Trp Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp
 115 120 125
Pro Arg His Arg Ser Arg Asn Val Gly Lys Val Ile Asp Thr Leu Thr
 130 135 140
Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Phe Arg Val Gly Ala Phe

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 145 150 155 160
Leu Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu
165 170 175
```

For all numeric identifiers in this file, please remove the space between the number and the right hand bracket, ">". Please add a space between the right hand bracket, ">", and the response. Using numeric identifier <160> as an example, change the incorrect format "<160 >8", to the correct format "<160> 8". Please make these changes to all the numeric identifiers found in this file.

Numeric identifiers "<130> File Reference" is mandatory on first submission. Please add numeric identifier <130> with the appropriate response. Note: for amended responses numeric identifiers "<140> Current Application Number" and numeric identifier "<141> Current Filing Date" are mandatory fields.

2.
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<210 >2
<211 >160
<212 >TRP
<213 >Hepatitis C virus
<400 >2
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For SEQ ID # 2, numeric identifier "<211> Type" can only be one of three choices PRT, DNA, or RNA. Please make all necessary changes.

****

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================================================================================
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Validation Report for Application: 09269897 cannot be generated.

This file is not a valid Sequence Listing File.

Timestamp: Tue Jul 28 11:03:27 EDT 2009
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================================================================================
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Met Lys Ala Ile Phe Val Leu Lys Gly Ser Leu Asp Arg Asp Pro Glu
5 10 15

Phe Met Gly Thr Asn Pro Lys Pro Gin Arg Lys Thr Lys Arg Asn Thr
20 25 30

Asn Arg Arg Pro Gin Asp Val Lys Phe Pro Gly Gly Gly Gin Ile Val
35 40 45

Gly Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg
50 55 60

Ala Thr Arg Lys Thr Ser Lys Arg Ser Gin Pro Arg Gly Gly Arg Arg
65 70 75 80

Pro Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro
85 90 95

Gly Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly
100 105 110

Trp Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp
115 120 125

Pro Arg His Arg Ser Arg Asn Val Gly Lys Val Ile Asp Thr Leu Thr
130 135 140

Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Phe Arg Val Gly Ala Phe
145 150 155 160

Leu Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu
165 170 175

Met Gly Thr Asn Pro Lys Pro Gin Arg Lys Thr Lys Arg Asn Thr Asn
5 10 15

Arg Arg Pro Gin Asp Val Lys Phe Pro Gly Gly Gly Gin Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Lys Arg Ser Gin Pro Arg Gly Gly Arg Arg Pro
50 55 60

Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser 'Prp Gly Pro Thr Asp Pro
100 105 110

Arg His Arg Ser Arg Asn Val Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Phe Arg Val Gly Ala Phe Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

<210 >3
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<212 >PRT
<213 >Artificial Sequence
<220>
<223 >
<400 >3
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Leu Pro Arg Arg
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<212 >PRT
<213 >Artificial Sequence
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<223 >
<400 >4
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5 10
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<211 >21
<212 >PRT
<213 >Artificial Sequence
<220>
<223 >
<400 >5
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Ser Arg Asn Val Gly
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<210 >6
<211 >20
<212 >PRT
<213 >Artificial Sequence
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<230 >
<400 >6
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1 5 10 15
Thr Cys Gly Phe
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<210 >7

<211 >24
<212 >DNA
<213 >Artificial Sequence
<220 >Probe
<230 >Synthetic DNA
<400 >7
gaattcatgg gcacgaatcc taaa 24
<210 >8
<211 >21
<212 >DNA
<213 >Artificial Sequence
<220 >Probe

<230 >Synthetic DNA
<400 >8
ttagtcctcc agaacccgga c 21